

Figure1

A. <http://zinc.docking.org/substances/search/?q=sinensetin>

B.

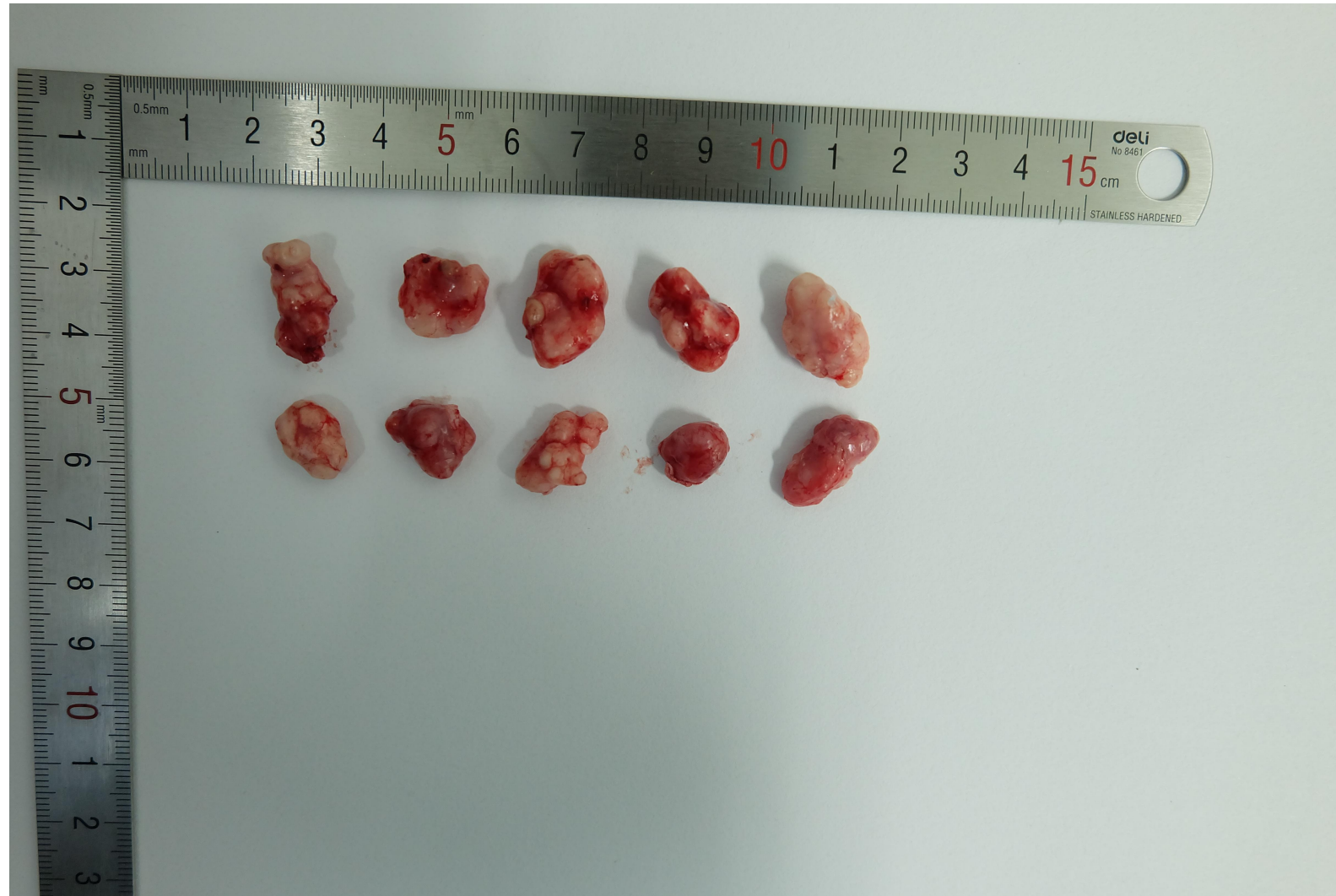


Figure1

D.CON

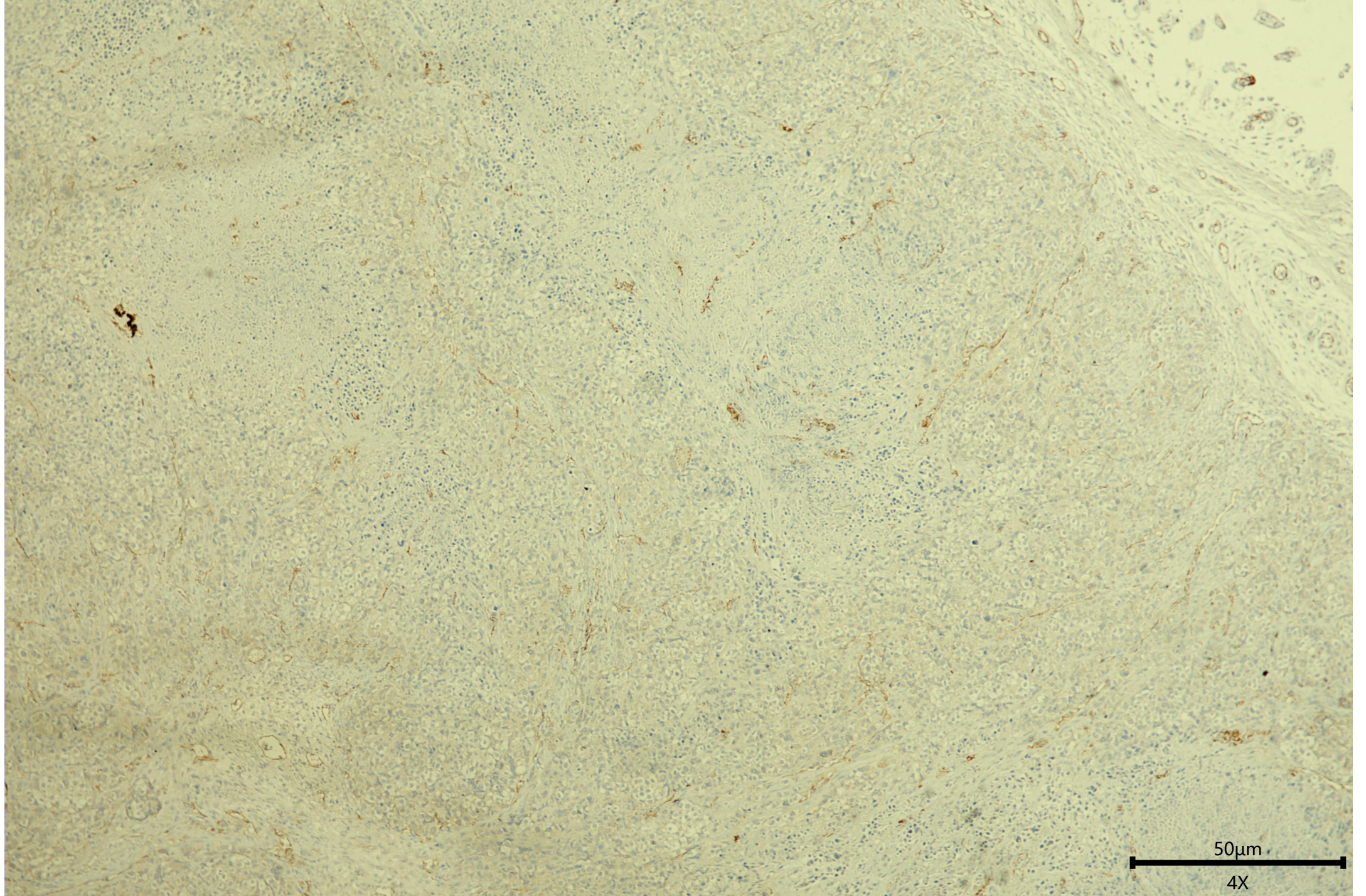


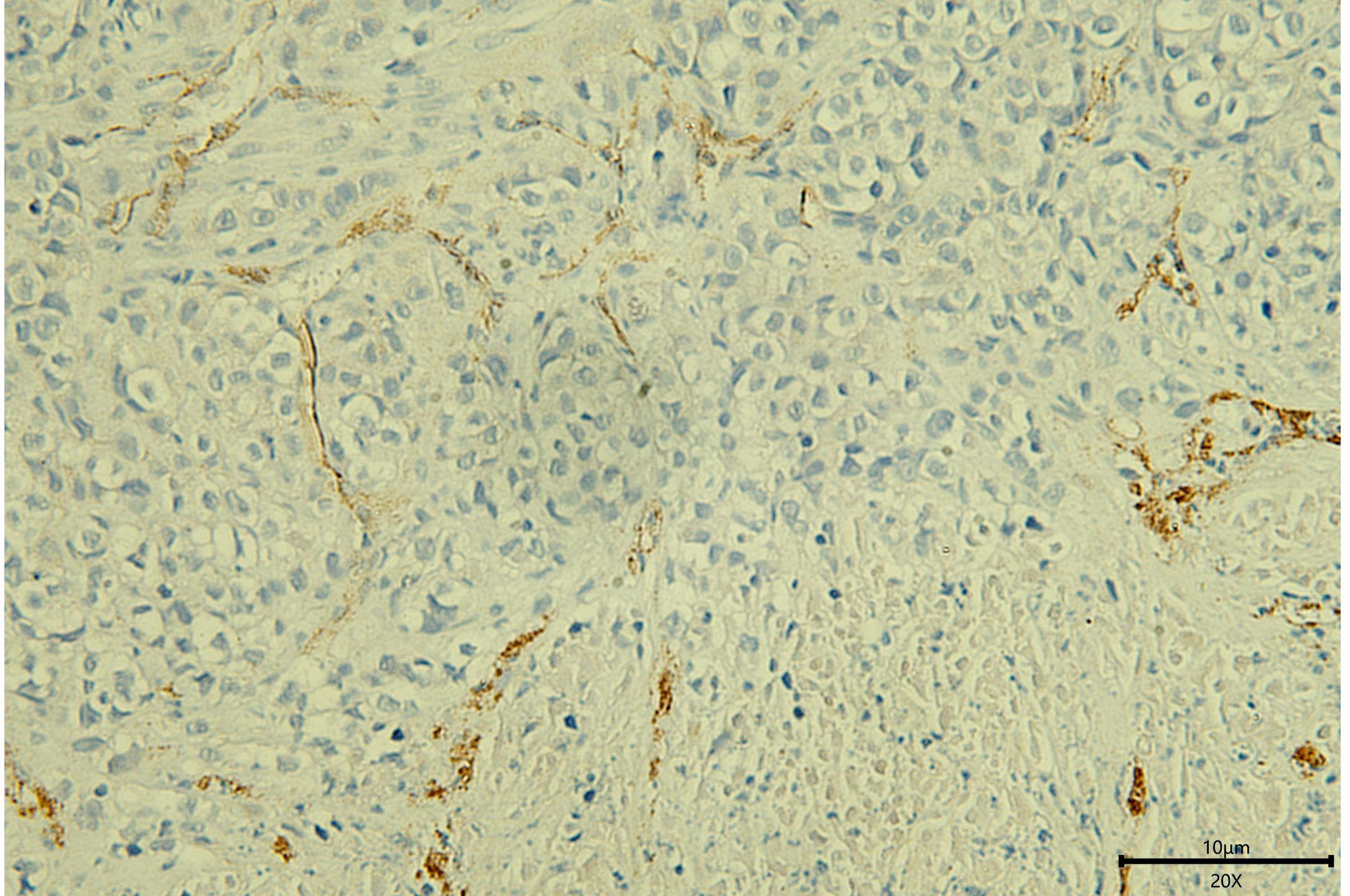
Figure1

D.SIN



Figure1

D.CON

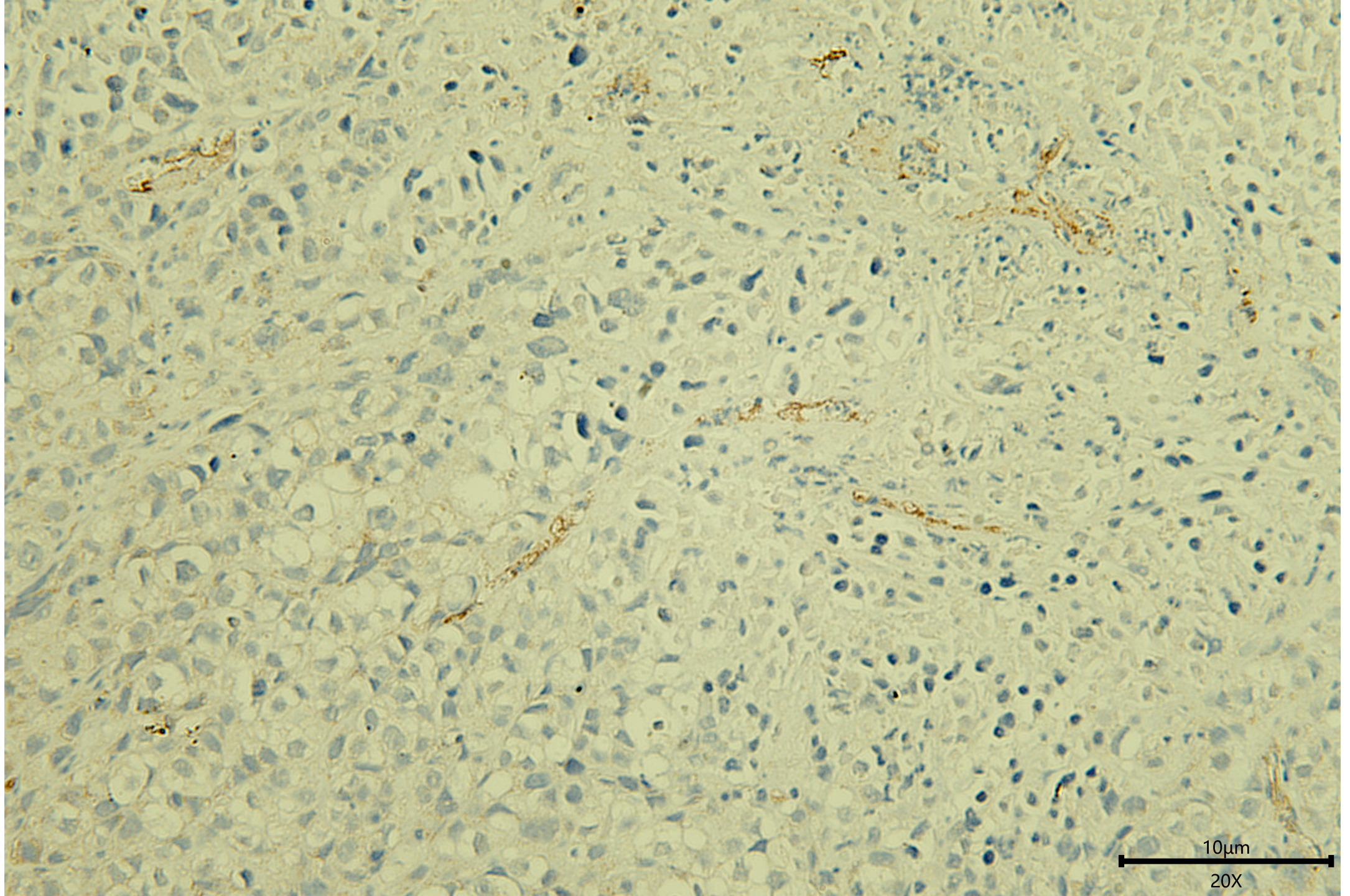


10μm

20X

Figure1

D.SIN

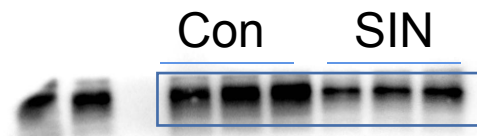


10μm

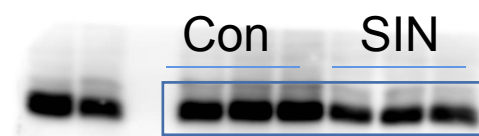
20X

Figure1

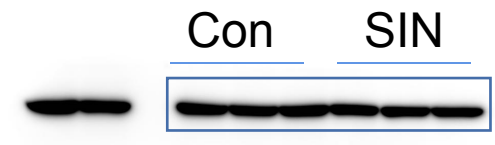
CD31



VEGF

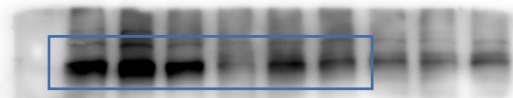


α -Tublin



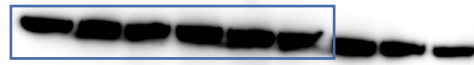
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Con SIN

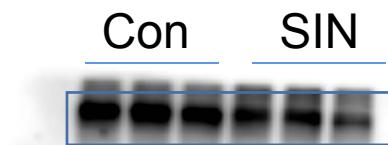


α -Tublin

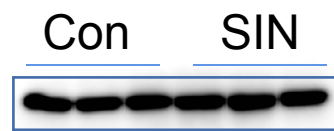
Con SIN



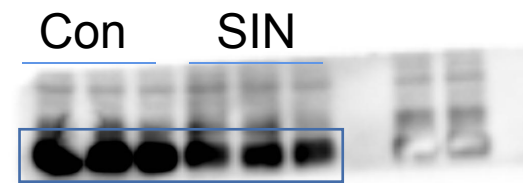
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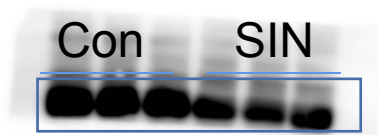
α -Tublin



VEGF



VEGF



α -Tublin

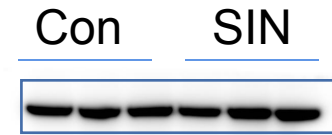
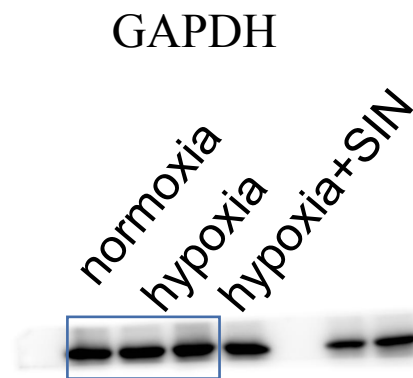
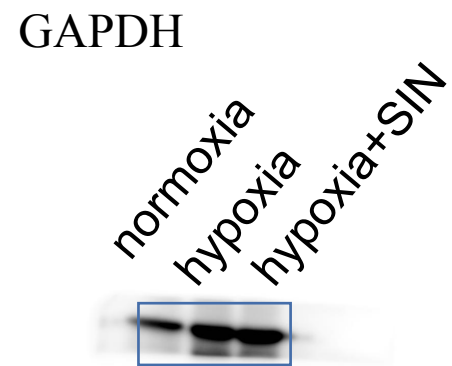
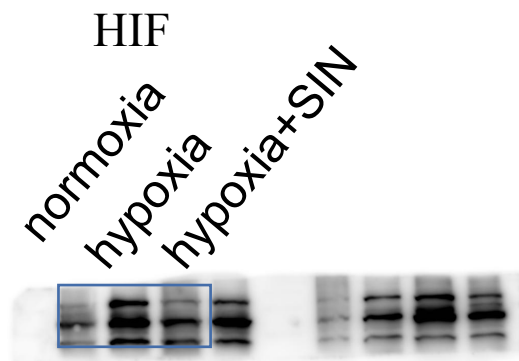
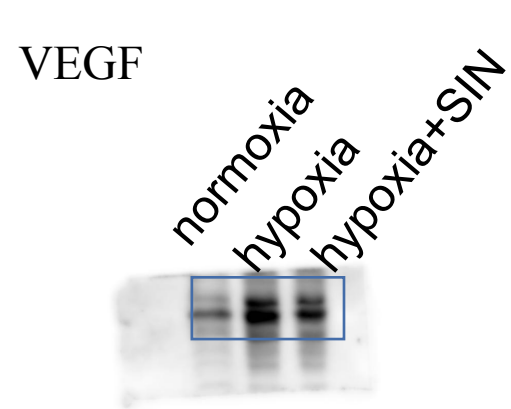


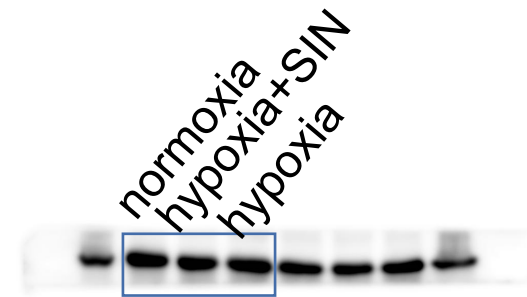
Figure 2A



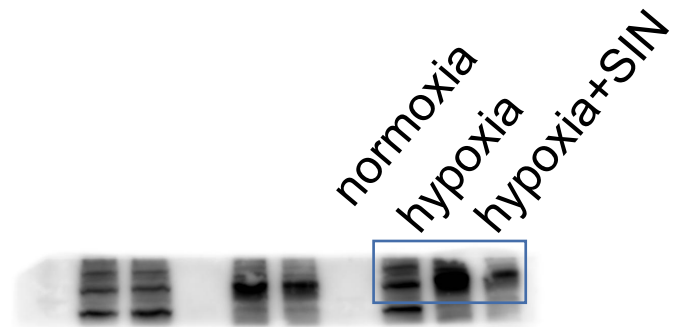
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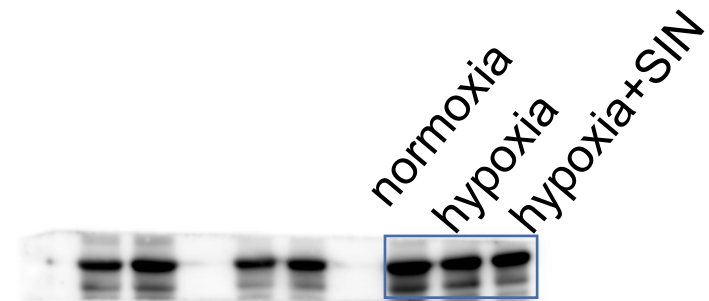
GAPDH



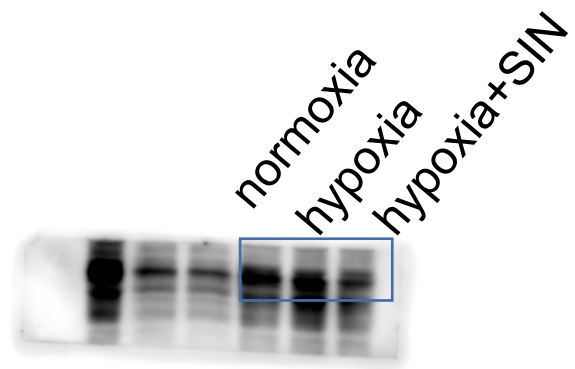
HIF



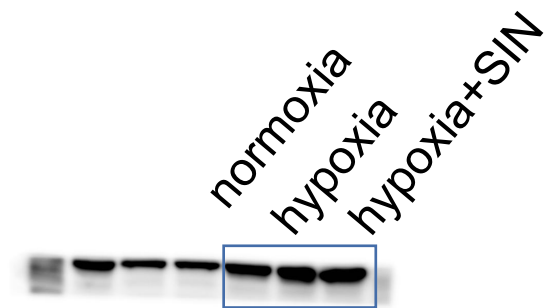
GAPDH



VEGF



GAPDH



VEGF

normoxia
hypoxia
hypoxia+SIN



GAPDH

normoxia
hypoxia
hypoxia+SIN



Figure2D

D.CON

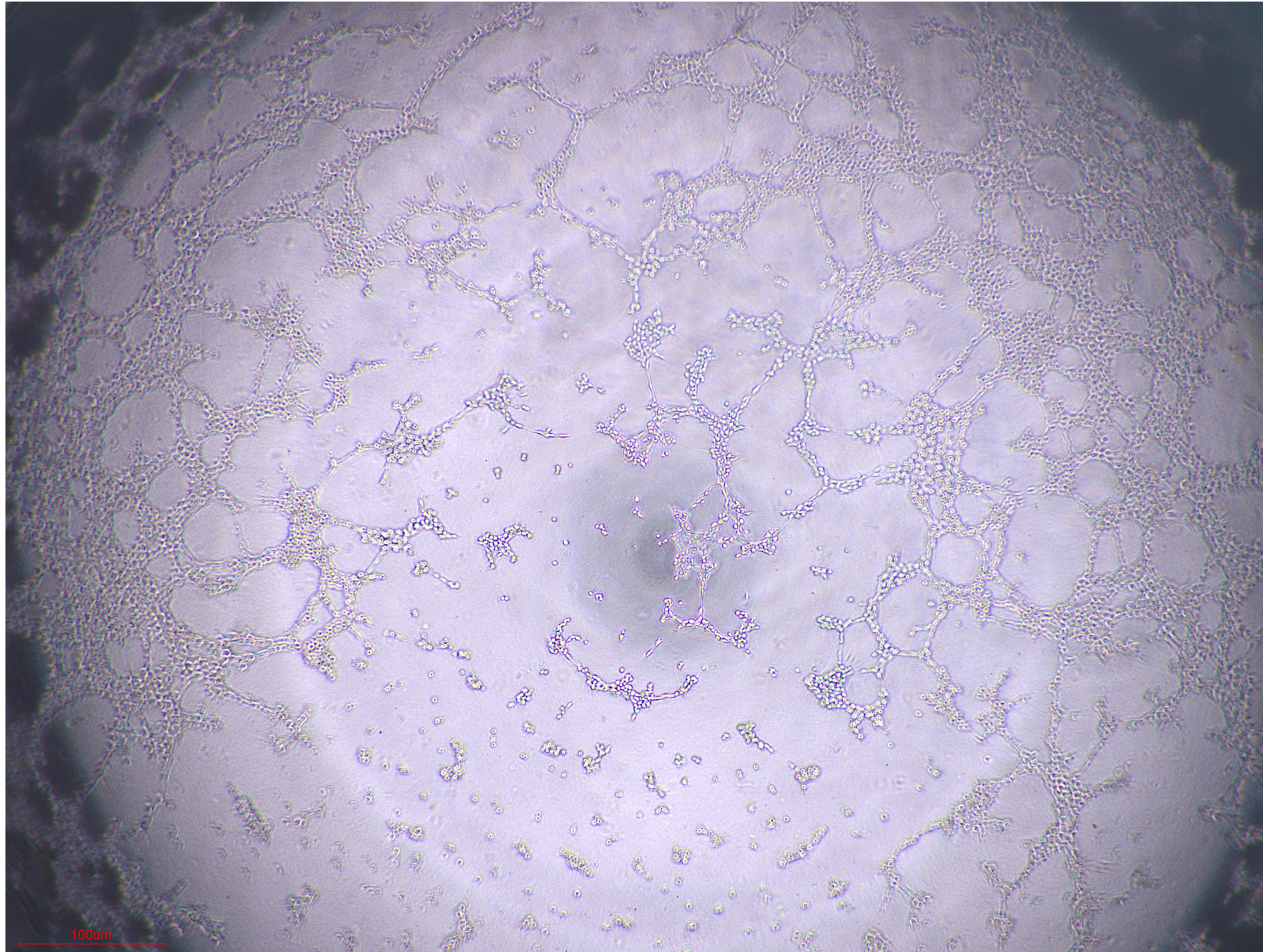


Figure2D

D.SIN

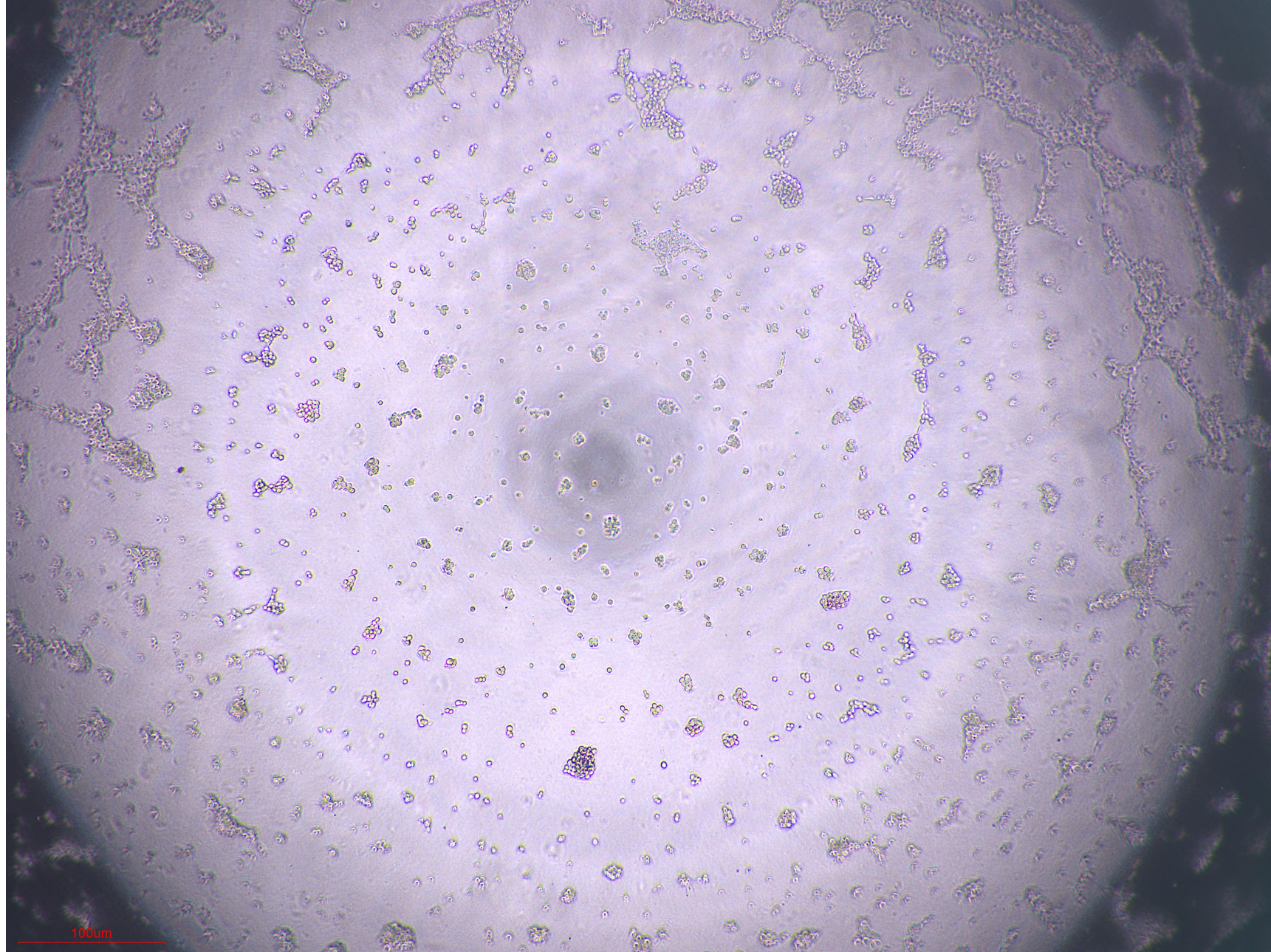


Figure3C

CON 0h

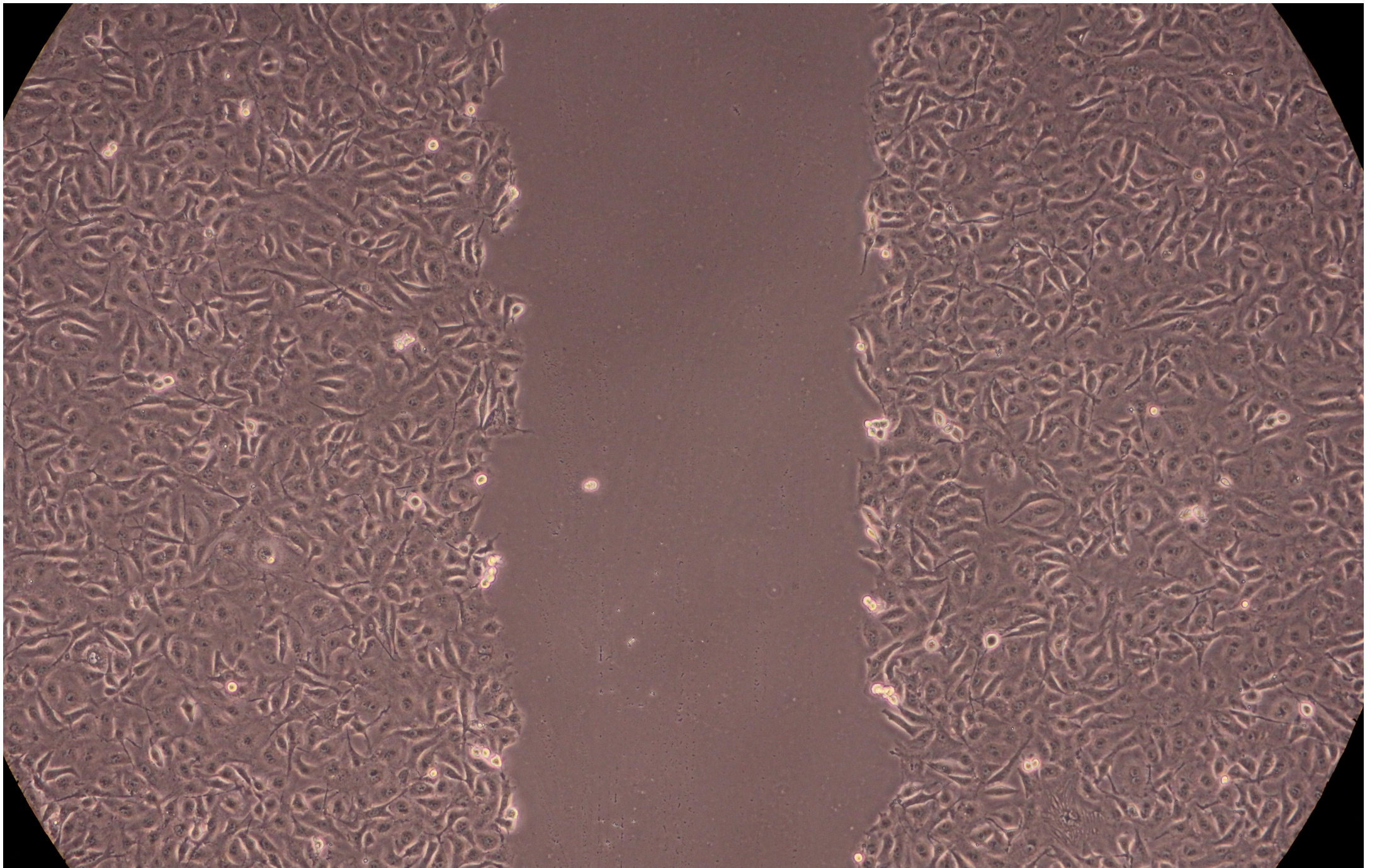


Figure3C

CON 24h

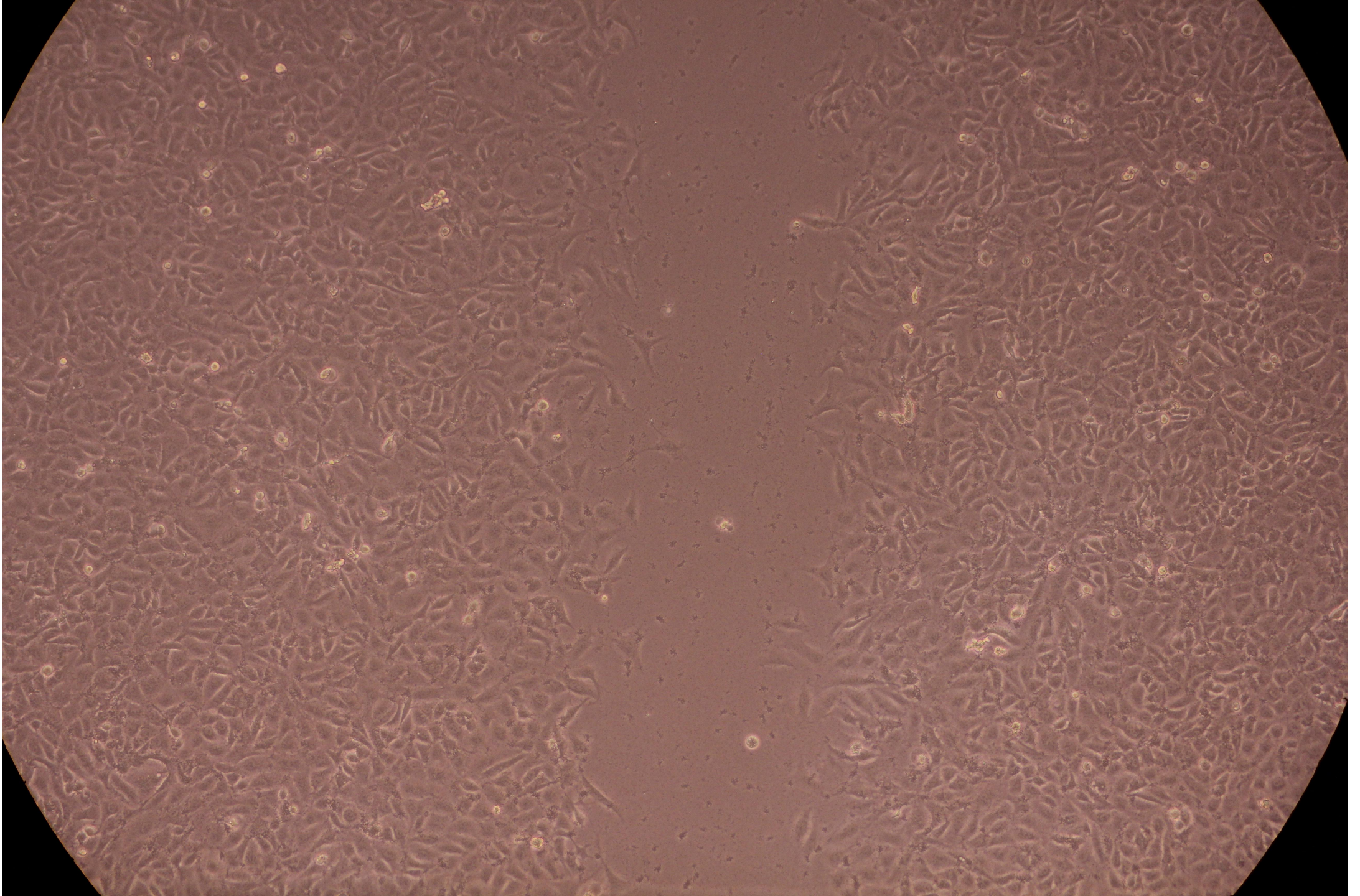


Figure 3C

VEGF
0h

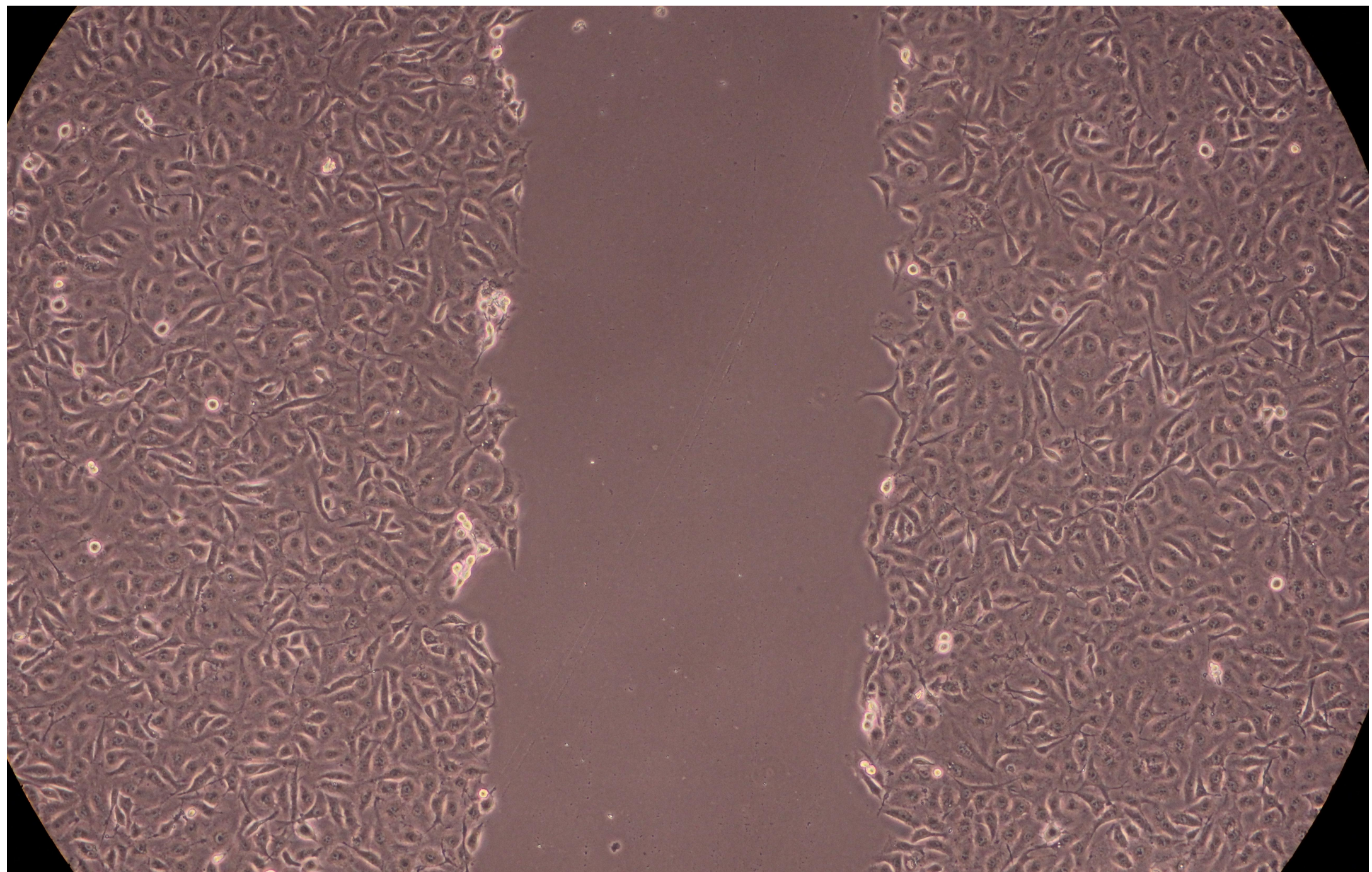


Figure 3C

VEGF
24h

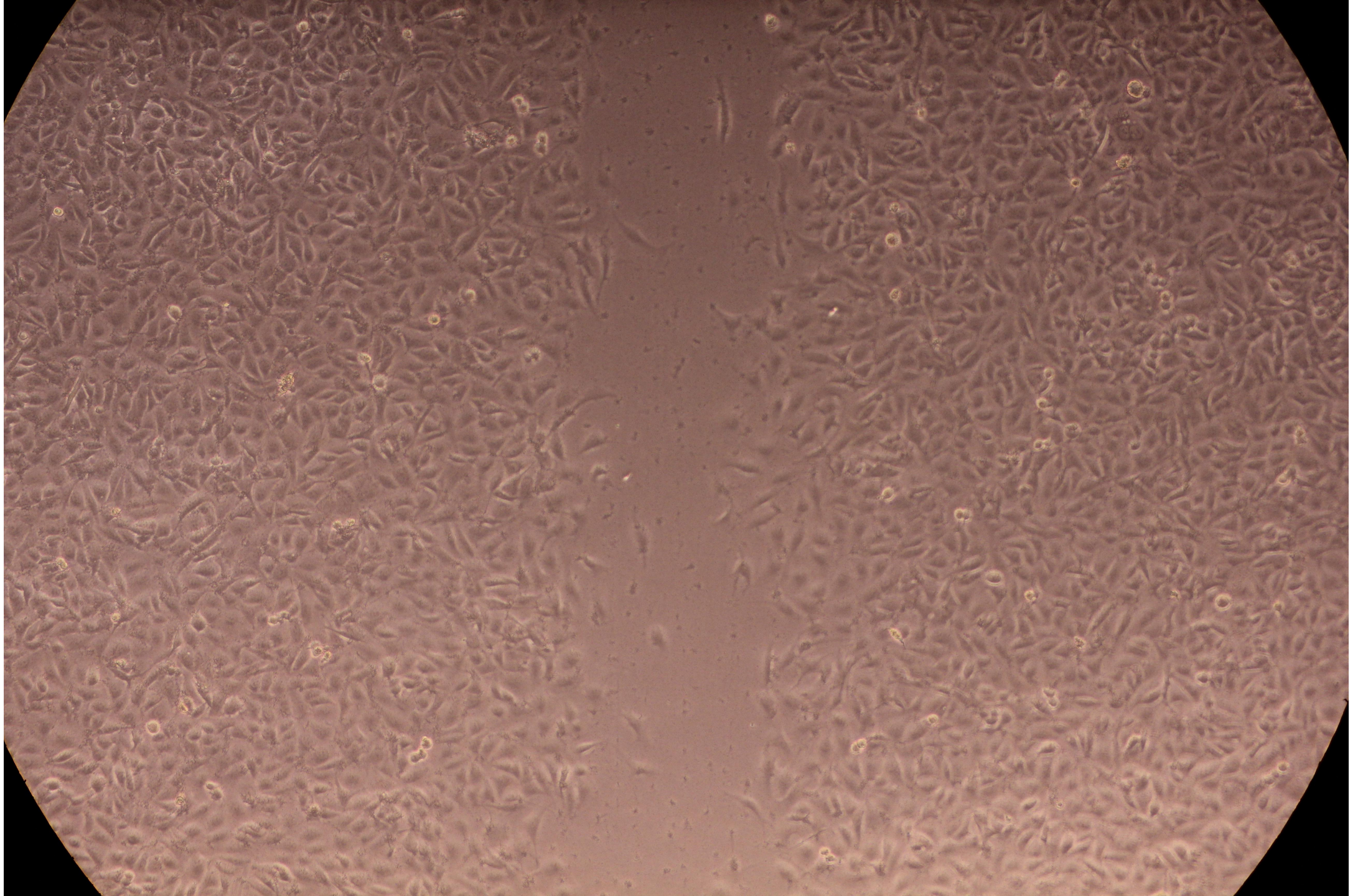


Figure3C

VEGF+
SIN 0h

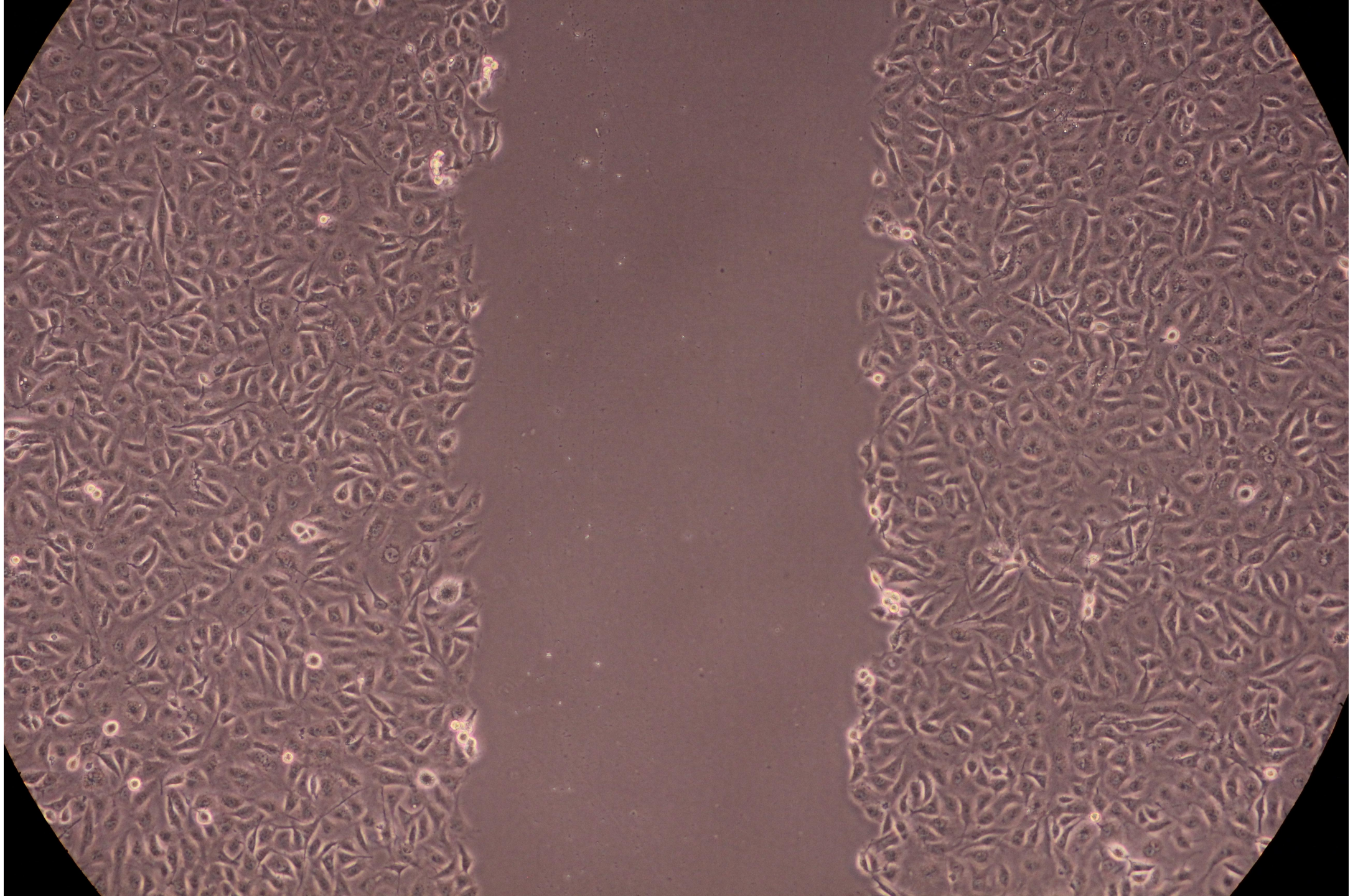


Figure 3C

VEGF+
SIN 24h

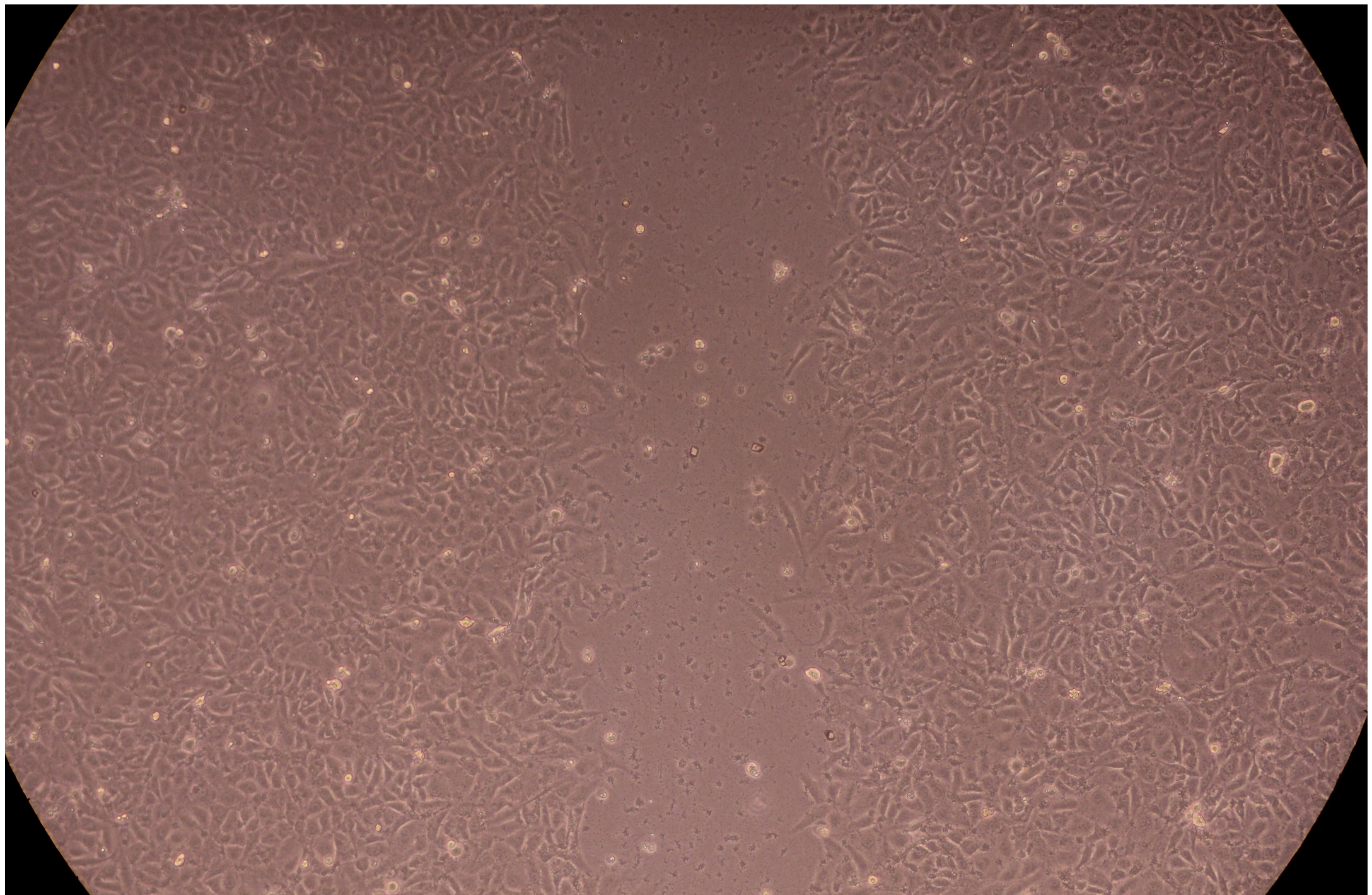


Figure3D

CON

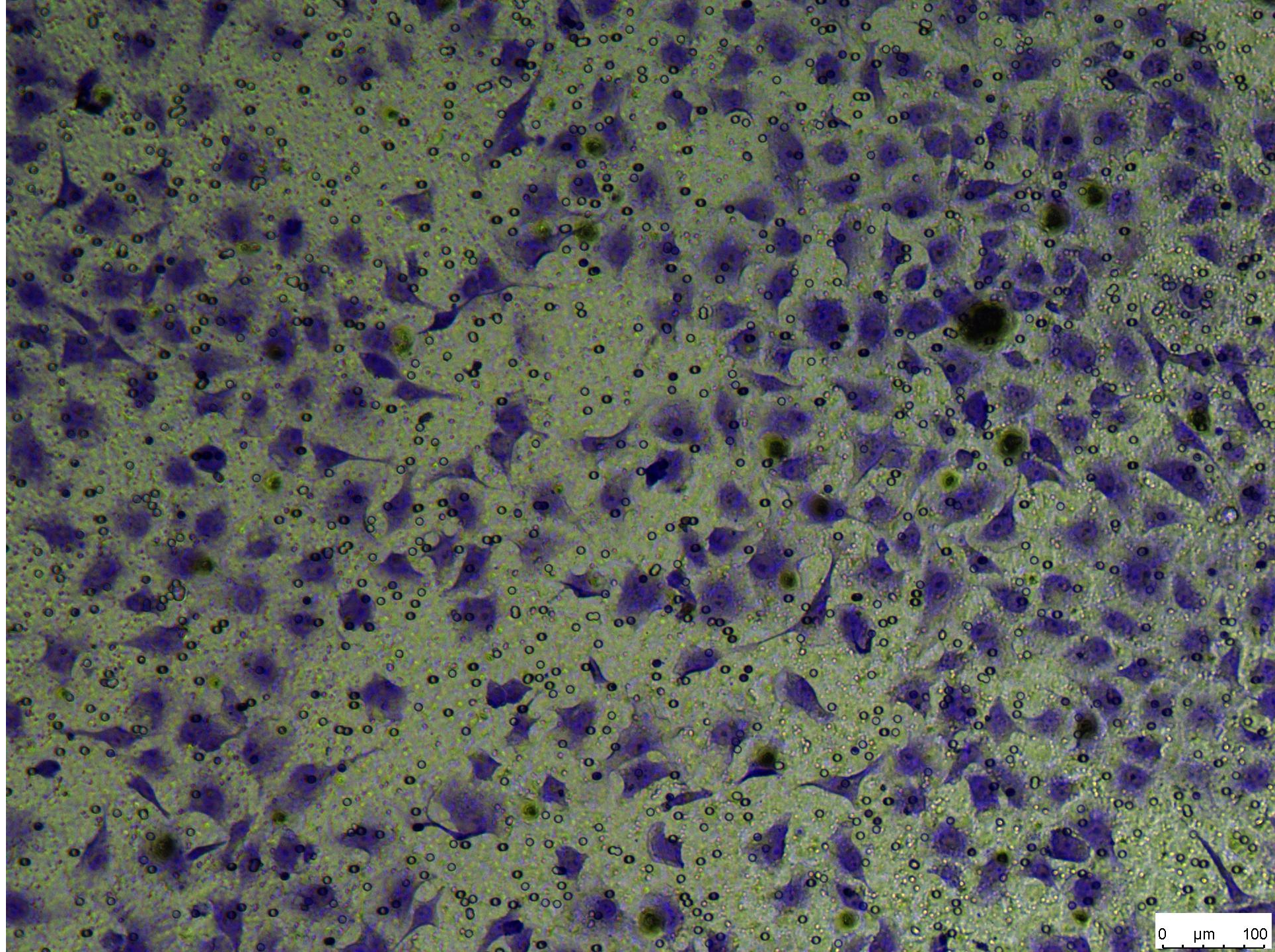


Figure3D

VEGF

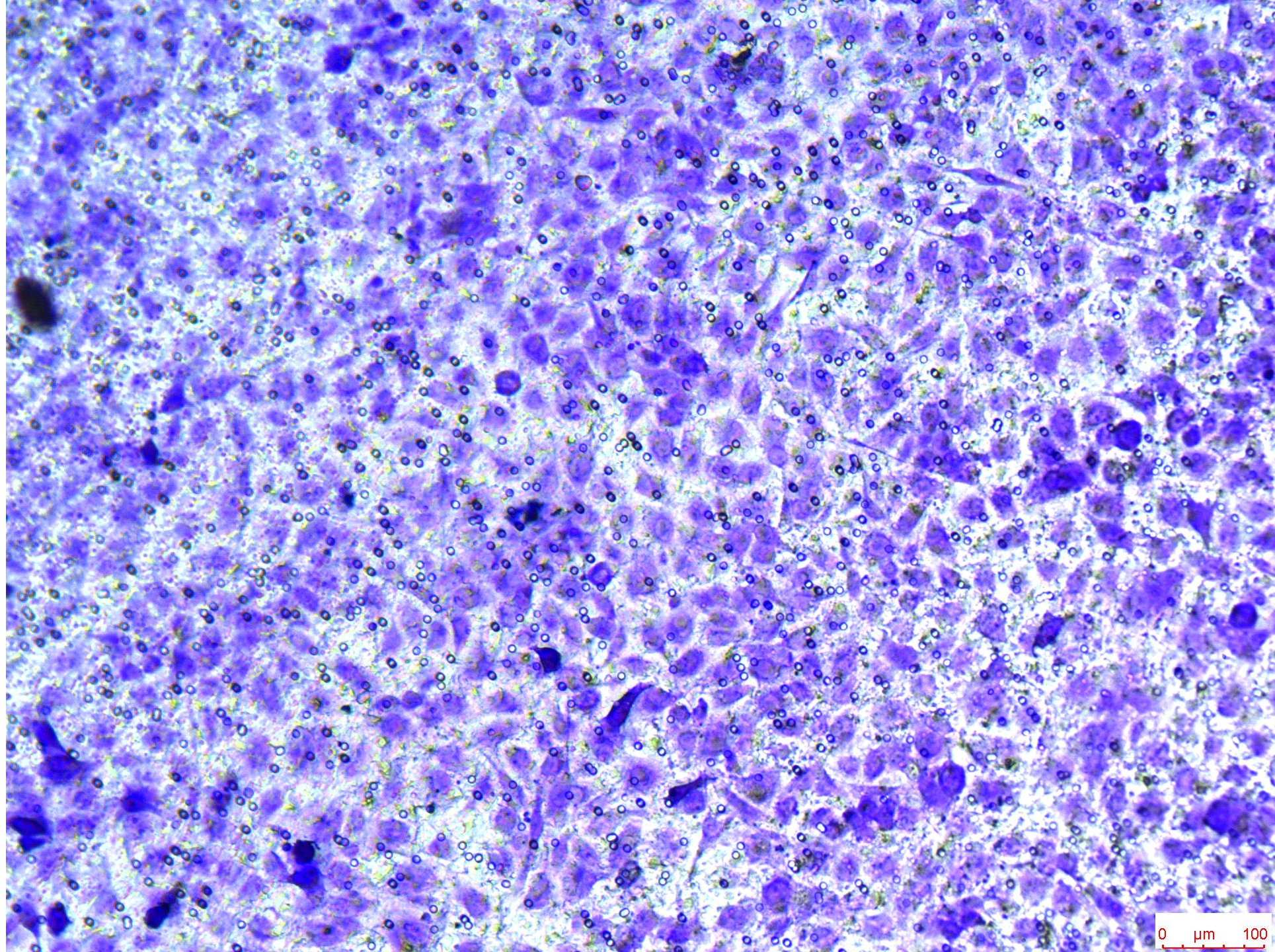


Figure3D

VEGF+S
IN

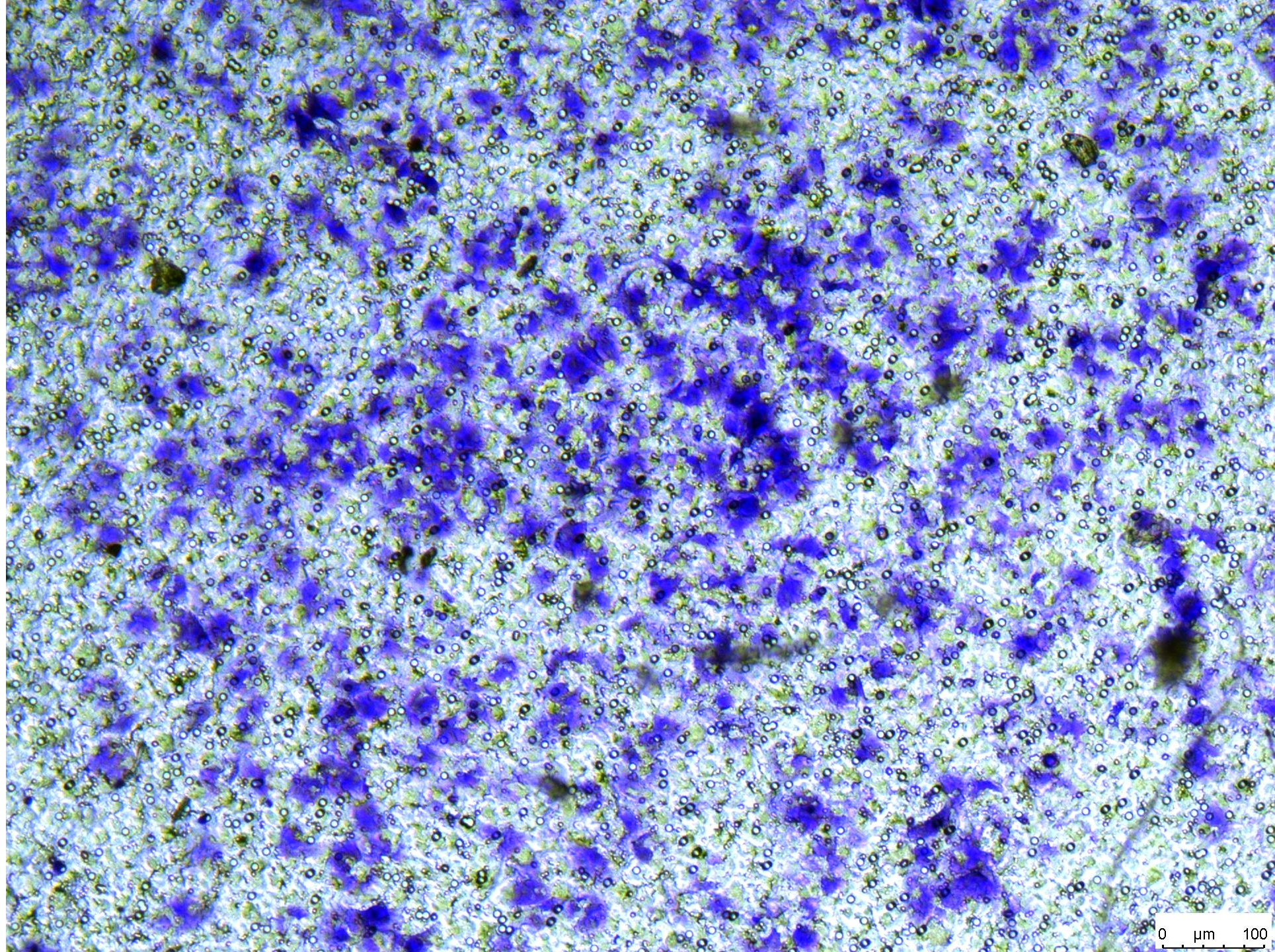


Figure3E

CON

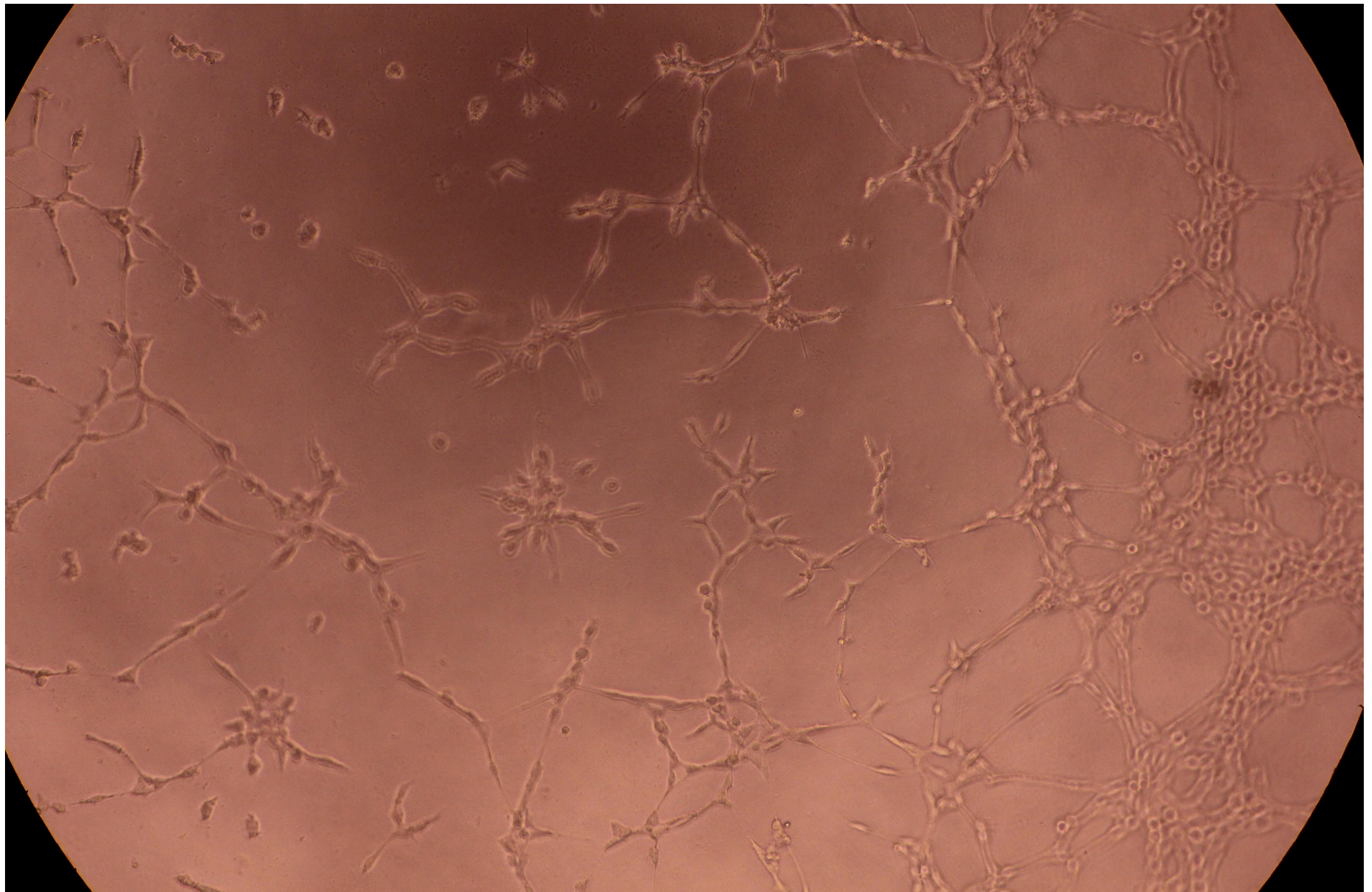


Figure 3E

VEGF

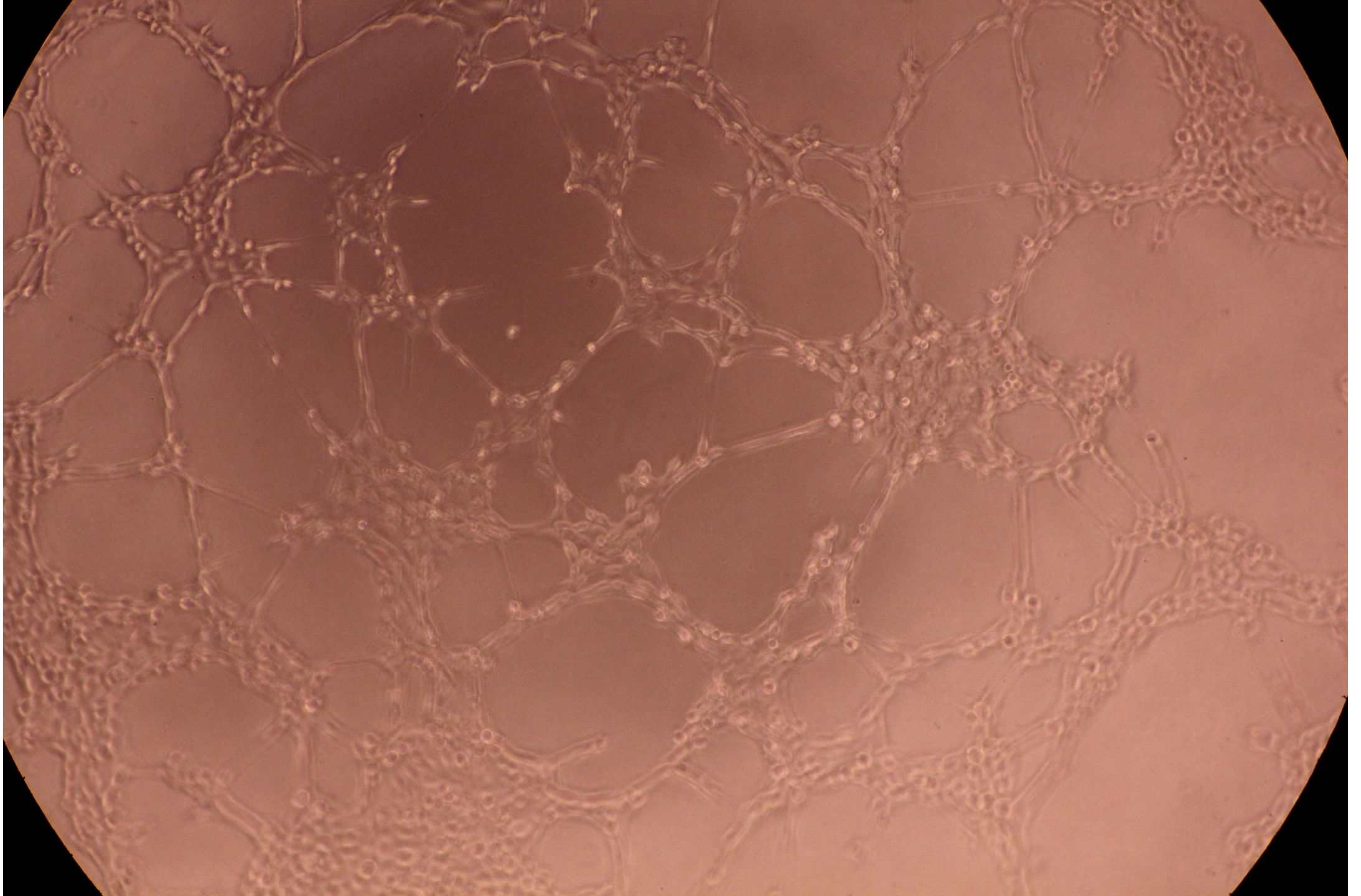


Figure3E

VEGF+SIN

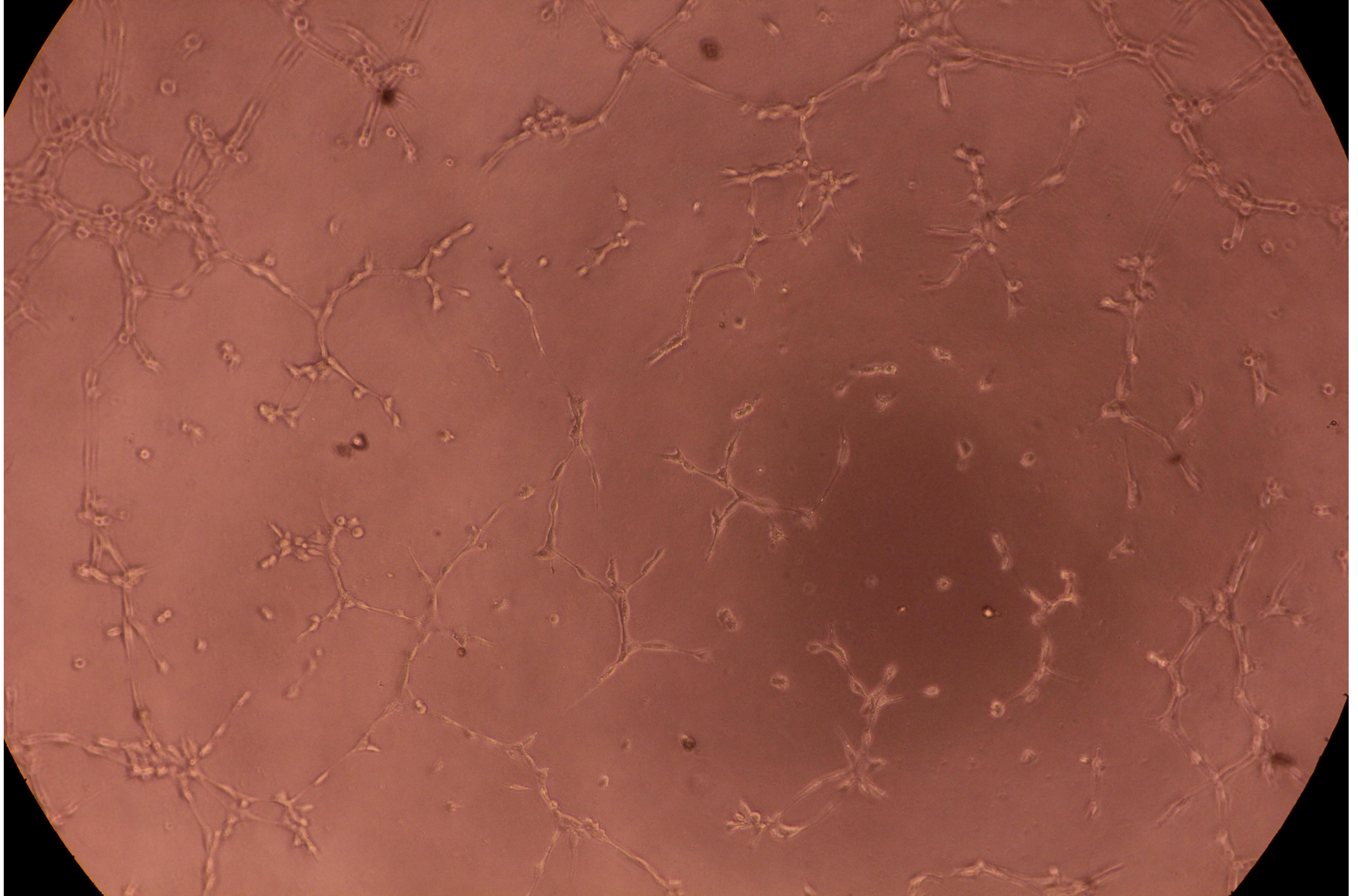
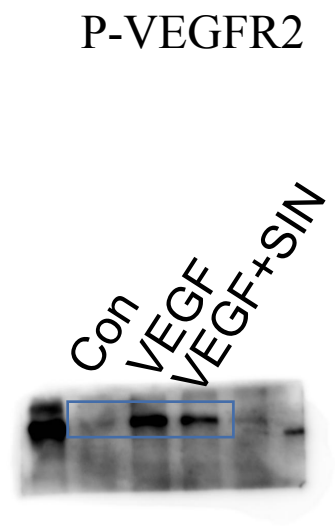


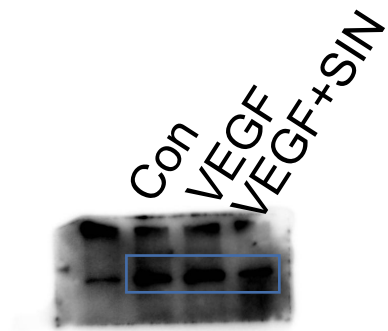
Figure4A



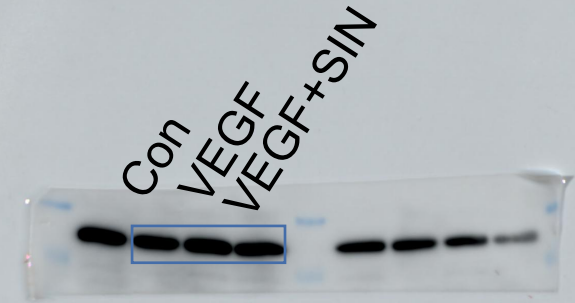
P-VEGFR2



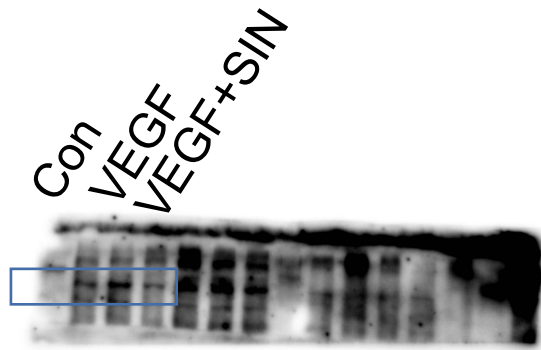
VEGFR2



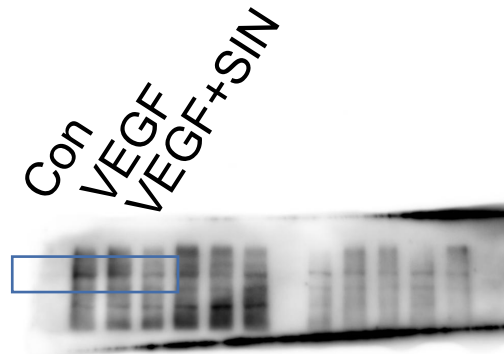
β -ACTIN



P-VEGFR2



VEGFR2



GAPDH

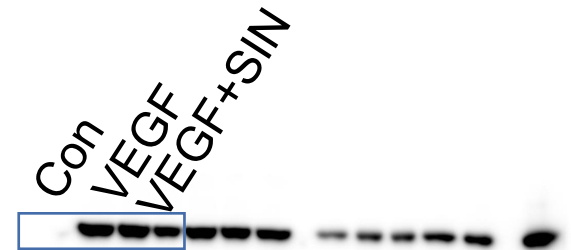
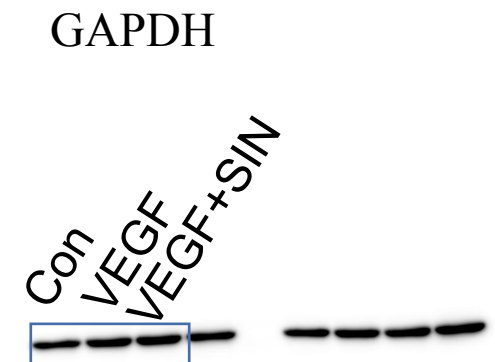
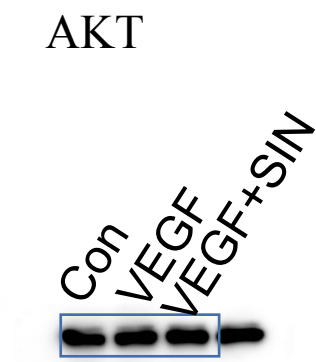
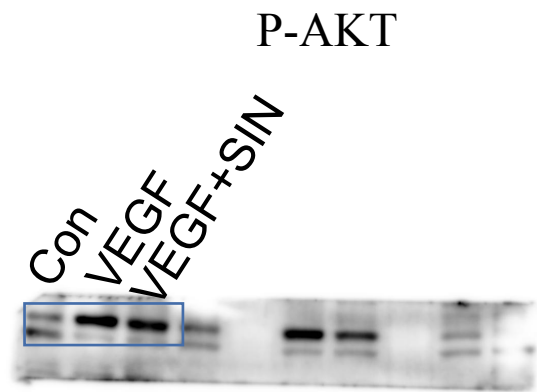


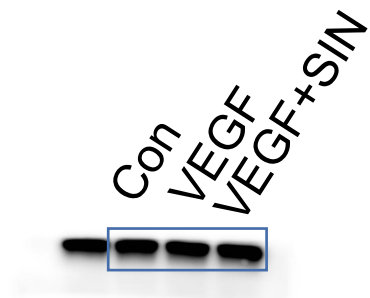
Figure4B



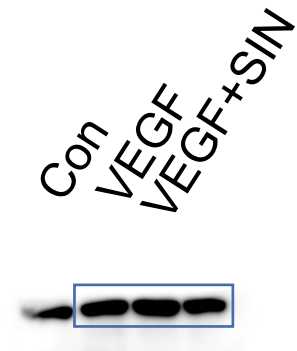
P-AKT



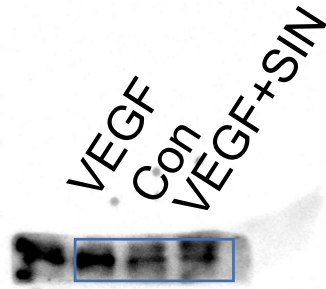
AKT



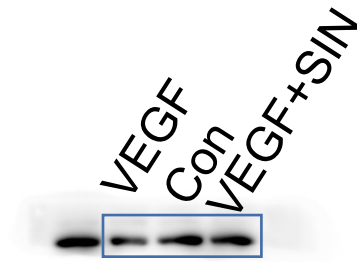
GAPDH



P-AKT



AKT



GAPDH

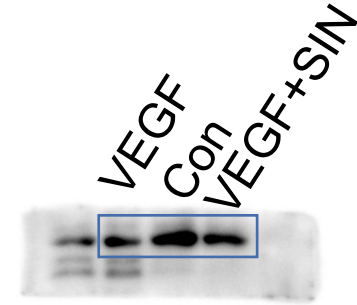
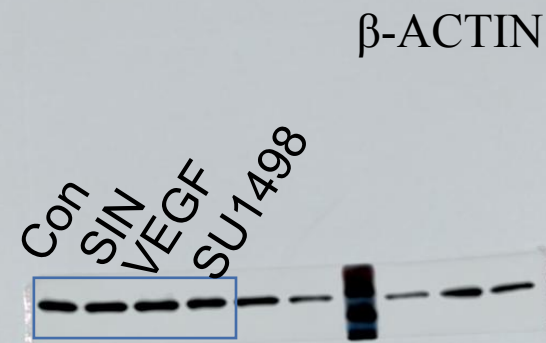
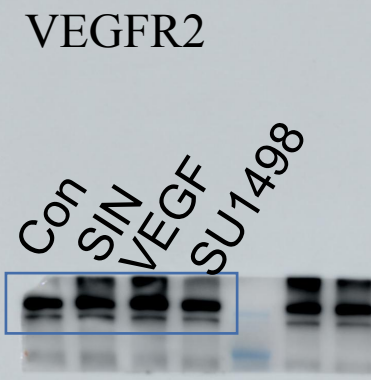
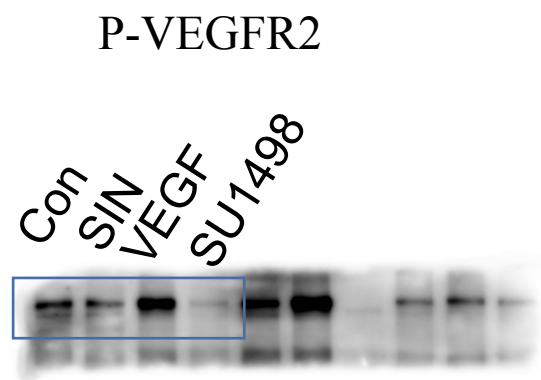
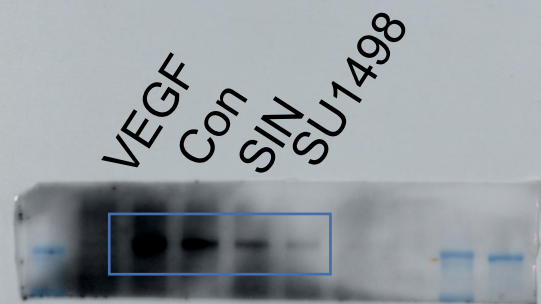


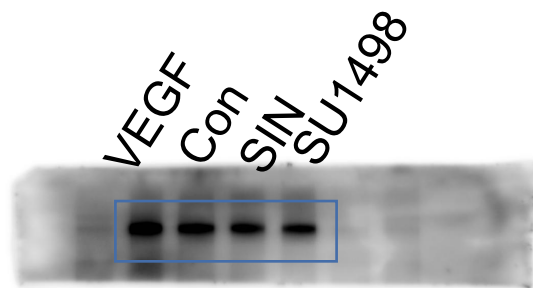
Figure 5A



P-VEGFR2



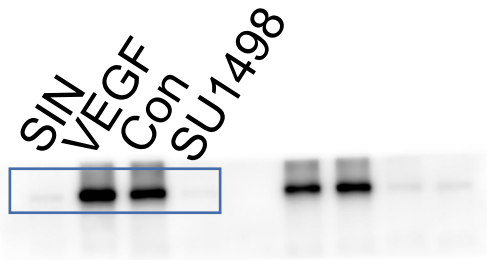
VEGFR2



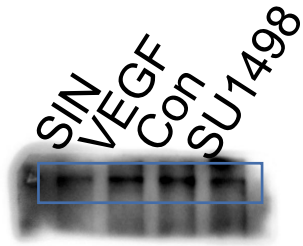
β -ACTIN



P-VEGFR2



VEGFR2

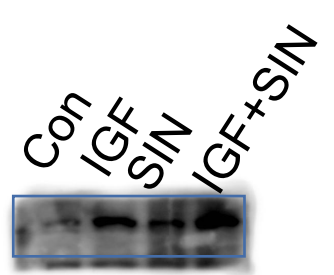


GAPDH

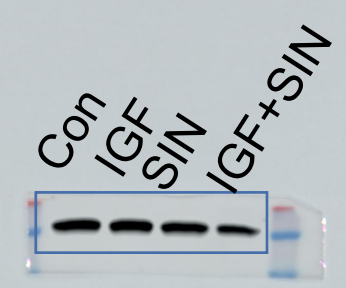


Figure 5B

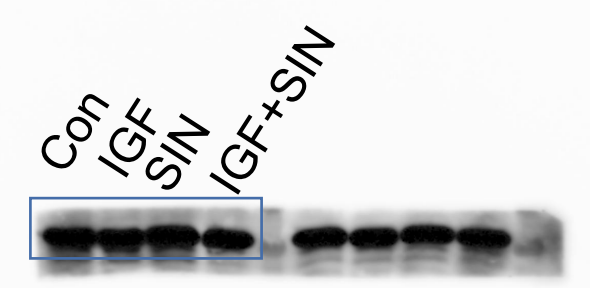
P-AKT



AKT



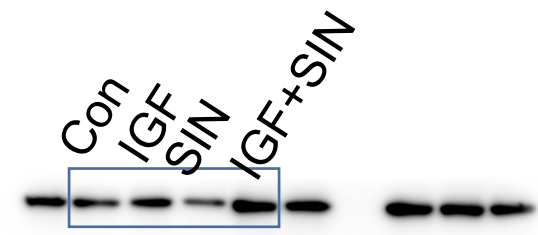
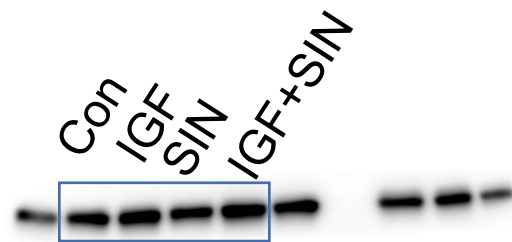
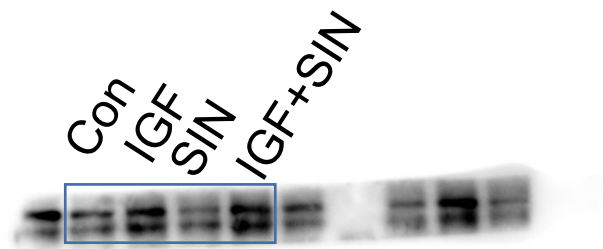
GAPDH



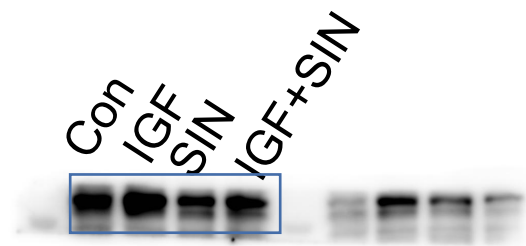
P-AKT

AKT

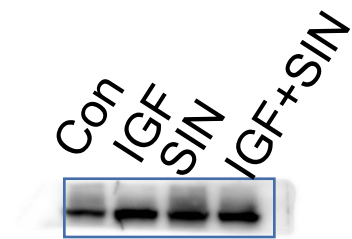
GAPDH



P-AKT



AKT



GAPDH

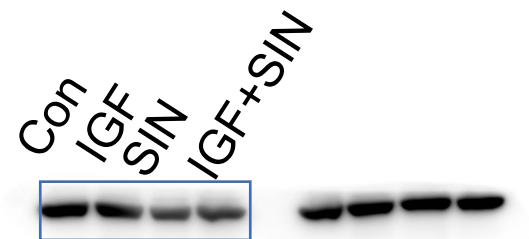


Figure6A



Figure6B

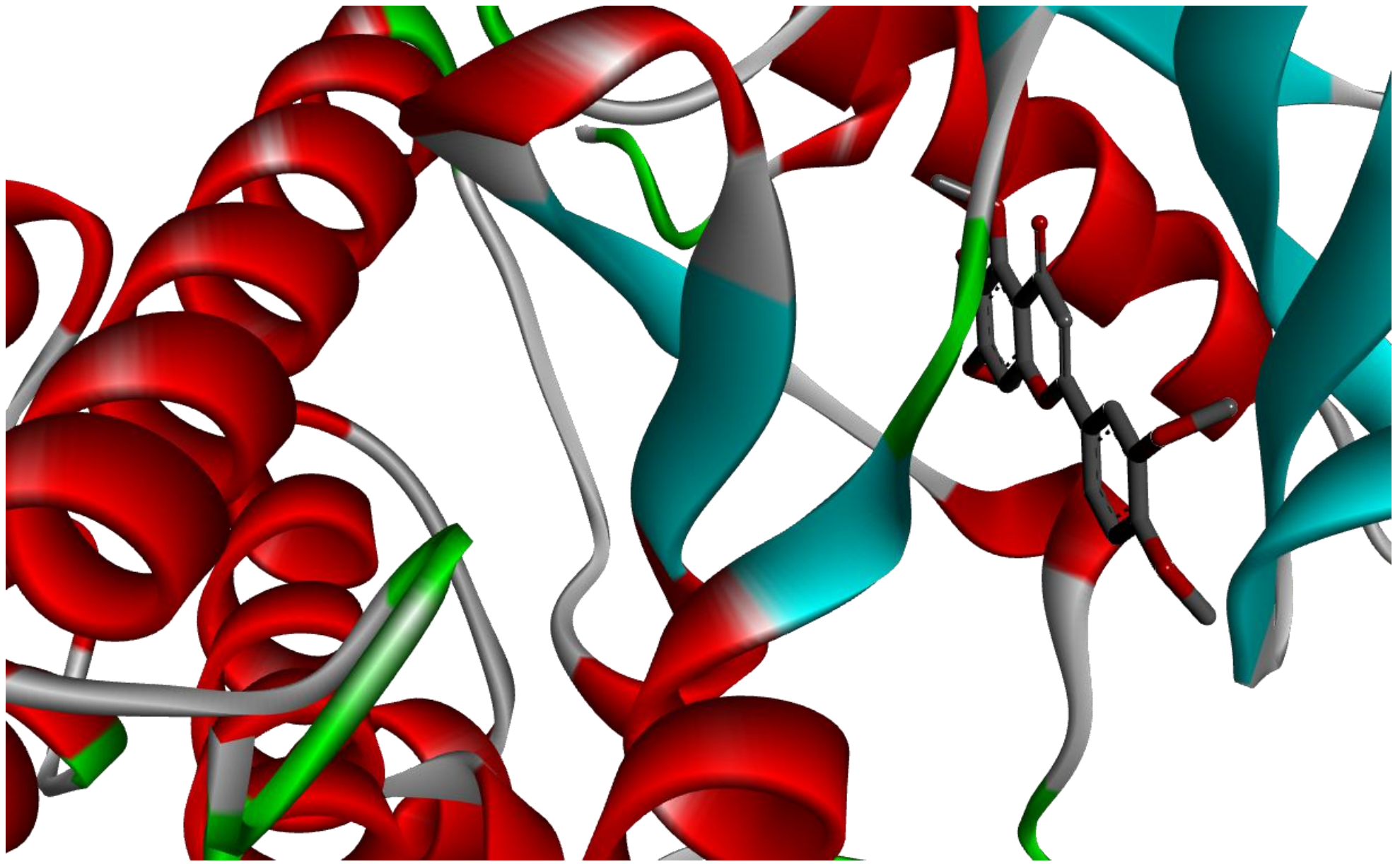
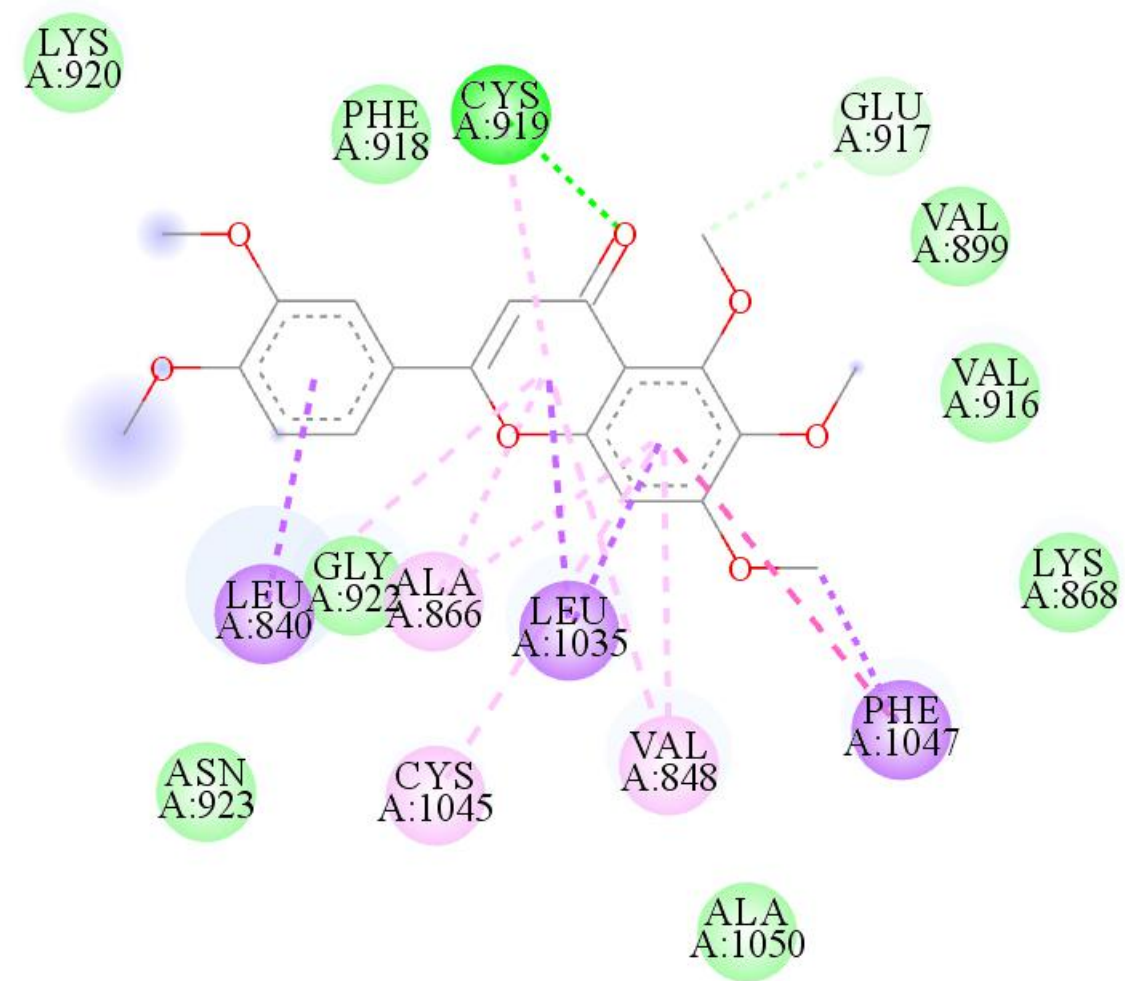


Figure 6C

A 3D molecular model showing a protein structure (represented by a semi-transparent surface) and a bound ligand (represented by a stick model). The protein surface is color-coded according to a legend on the left. The legend is a vertical color bar with the following labels from top to bottom: "Hydrophobic" (grey), "Donor" (pink), "Acceptor" (green), and "Hydrophilic" (white). The protein surface shows green regions (Acceptor) and pink regions (Donor). The ligand is a small molecule with a central ring and several side chains, colored in grey (carbon), red (oxygen), and white (hydrogen).

Figure6D



Interactions

- van der Waals
- Conventional Hydrogen Bond
- Carbon Hydrogen Bond

- Pi-Sigma
- Pi-Pi T-shaped
- Pi-Alkyl